

FIGURE 1

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A. The amino terminal sequence of IL13-gC

B. The sequence of the gB_{Δpoly(K)} domain

GGGTCCCTGGTGGCGCTGGCGGCTCGGAGTTCCCCCGGACGGCTGGGTGCGGGCGCGACCCAGGCGGC
GAACGGGGGACCTGCCACTCCGGGCGGCCGGCCCTGGCCCCGGCCAAAGGGGGATCCAAACCGAAG
AAAGAACAGAAAACCGAAAACCCCCAAAGCGCGCGCGCGCGACAAACCGGACCGTGC CGCGGGCCA
CGCCACCCCTGCGCGAGCACCTGCGGGACATCAAGGGAGAACACCGATCAAACTTTACGTGTGCCCA
CCCCCCCACGGGCGCCACGGTGGTGCAGTTGAGCAGCCGCGCCGCTGCCCCGACCCGGGGAGGGTCAAGA

C. The amino terminal sequence of IL13-gD

FIGURE 2

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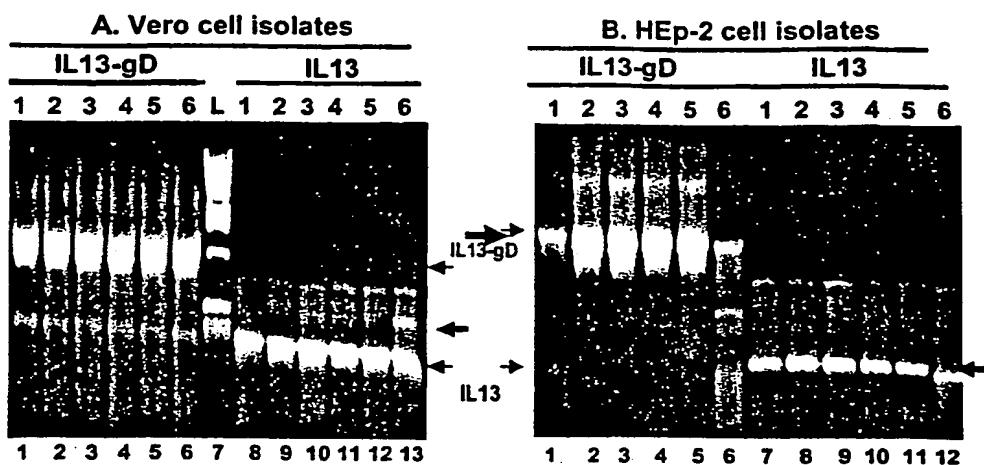
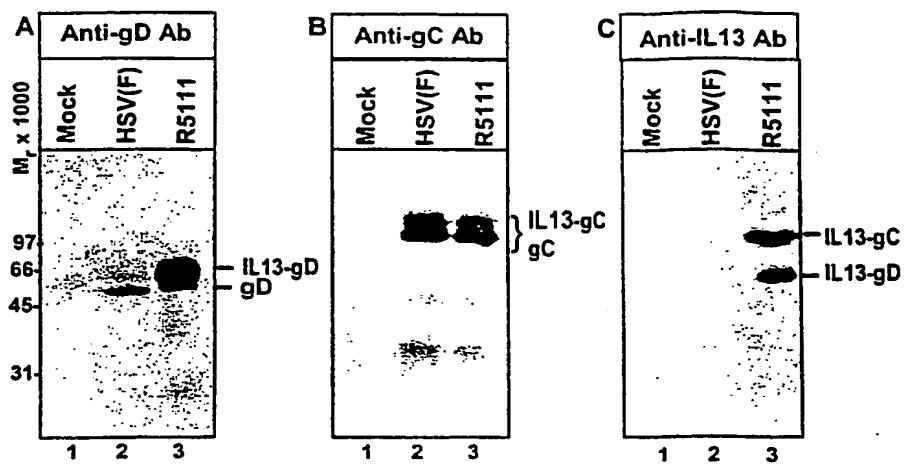


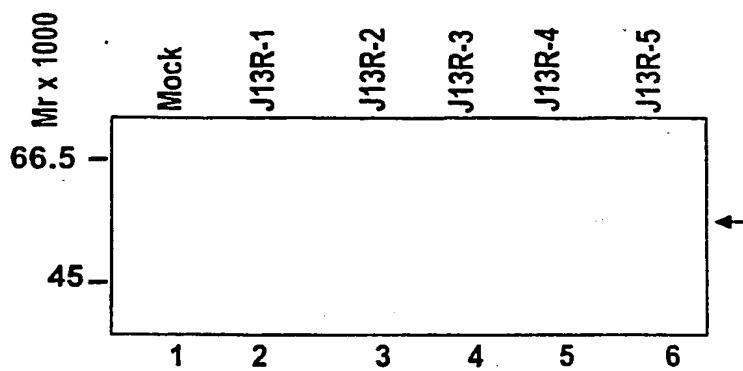
FIGURE 3

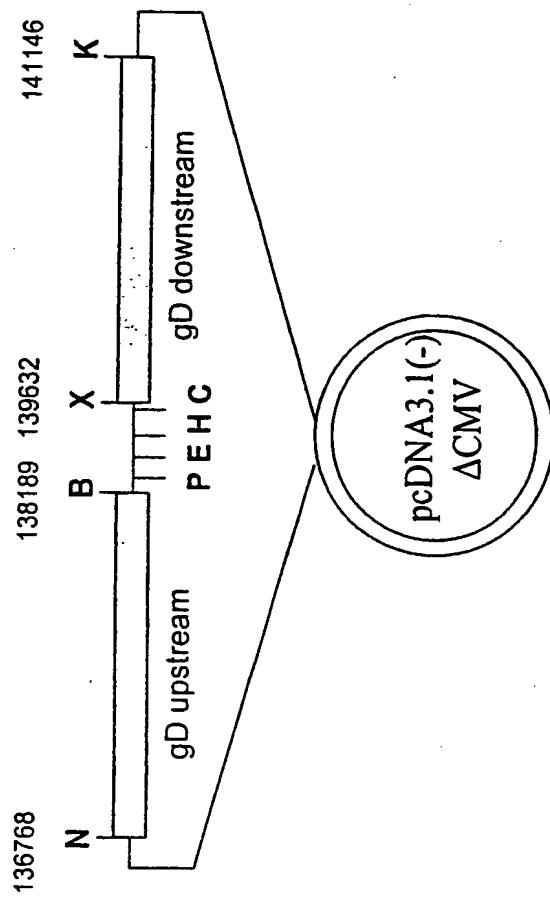
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**FIGURE 4****BEST AVAILABLE COPY**

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**FIGURE 5****BEST AVAILABLE COPY**



pgD- in pcDNA3.1(-) with CMV promoter collapsed by NruI/NheI digestion. The fragment containing gD upstream at NotI/BamHI, and gD downstream at Xhol/KpnI in pBluescript II SK was excised by NotI/KpnI and ligated into pcDNA3.1(-) in which the CMV promoter had been collapsed. N, NotI, B, BamHI, P, PstI, E, EcoRI, H, HindIII, C, Clal, X, Xhol, and K, KpnI.

FIGURE 6**BEST AVAILABLE COPY**